FORM PTO-1390 U.S.D.	DEPARTMENT OF COMMERCE PATENT AND TRADEMARK OFFI	ICE ATTORNEY'S DOCKET NUMBER				
(REV 10-94)	TO THE UNITED STATES	270.63USWO				
•	ED OFFICE (DO/EO/US)					
	NG UNDER 35 U.S.C. 371					
		US APPLICATION NO (If known, see 37 C F R 1 5)				
		Unknown <b>10</b> /049699				
INTERNATIONAL APPLICATION NO.	INTERNATIONAL FILING DATE	PRIORITY DATE CLAIMED				
PCT/CA00/01005	August 31, 2000	August 31, 1999				
TITLE OF INVENTION	<u> </u>					
METHOD OF CALIBRATING A SPECTRO	OSCOPIC DEVICE					
APPLICANT(S) FOR DO/EO/US						
CADELL						
Applicant herewith submits to the United States D	Designated/Elected Office (DO/EO/US) the followed	ing items and other information:				
1. [X] This is a <b>FIRST</b> submission of items con	ncerning a filing under 35 U.S.C. 371.					
2. [ ] This is a SECOND or SUBSEQUE	ENT submission of items concerning a filing under camination procedures (35 U.S.C. 371(f)) at any til	r 35 U.S.C. 371.				
examination until the expiration of the a	pplicable time limit set in 35 U.S.C. 371(b) and PC	CT Articles 22 and 39(1).				
4. [X] A proper Demand for International Preli	minary Examination was made by the 19th month	from the earliest claimed priority date.				
5. [X] A copy of the International Application a						
a. [X] is transmitted herewith (require     b. [X] has been transmitted by the Into	ed only if not transmitted by the International Bure ernational Bureau.	au).				
c. [ ] is not required, as the appl	ication was filed in the United States Receiving Of	ffice (RO/US)				
6. [ ] A translation of the International Ap	opplication into English (35 U.S.C 371(c)(2)).					
7. [X] Amendments to the claims of the Interna	(F)					
a. [ ] are transmitted herewith (r b. [ ] have been transmitted by the	equired only if not transmitted by the International he International Bureau.	l Bureau).				
<ul> <li>c. [ ] have not been made; howe</li> </ul>	ever, the time limit for making such amendments h	as NOT expired.				
d. [X] have not been made and will no	ot be made.					
8. [ ] A translation of the amendments to	the claims under PCT Article 19 (35 U.S.C. 371(c)	)(3)).				
	he inventor(s) (35 U.S.C. 371 (c)(4)).					
10. [ ] A translation of the annexes to the It (35 U.S.C. 371(c)(5)).	nternational Preliminary Examination Report unde	er PCT Article 36				
Items 11. to 16. below concern document(s) or it 11. [X] An Information Disclosure Statemen	information included: at under 37 CFR 1.97 and 1.98; form 1449; cited re	eferences.				
	ing. A separate cover sheet in compliance with 37					
13. [X] A FIRST preliminary amendment.						
[ ] A SECOND of SUBSEQUENT prel	liminary amendment					
14. [ ] A substitute specification.						
15. [ ] A change of power of attorney and/o	or address letter.					
16. [X] Other items or information: Application PCT/IPEA/402; PCT/IB/304; PCT/IB/308; PCT R	Data Sheet; International Preliminary Examination equest	n Report; International Search Report;				

JC11 Rec'd PCT/PTO 1 5 FEB 2002

				ATTÖRNEY'S DOCKET NUMBER		
Us. Application no (If known, see 37 C F R 15) Unknown 10/049699  INTERNATIONAL APPLICATION NO PCT/CA00/01005				270.63USWO		
17. [X] The following fees are submitted:				CALCULATIONS P	TO USE ONLY	
BASIC NATIONAL FEE (37 CFR 1.492(a) (1)-(5)): Search Report has been prepared by the EPO or JPO\$890.00					·	
	inary examination fee paid  1))		\$710.00			
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	al preliminary examination f n fee (37 CFR 1.445(a)(3)) p		\$1040.00			
	inary examination fee paid fied provisions of PCT Artic		\$100.00			
	ENTER APPROP	PRIATE BASIC FEE	AMOUNT =	\$890.00		
Surcharge of \$130.00 for furnishing the oath or declaration later than [] 20 [] 30 months from the earliest claimed priority date (37 CFR 1.492(e)).			\$0			
CLAIMS	NUMBER FILED	NUMBER EXTRA	RATE			
Total claims	-20 =	0	X \$18.00	\$0		
Independent claims	1 -3 =	0	X \$84.00	\$0		
MULTIPLE DEPENDE	NT CLAIM(S) (If applicable	le)	+ \$260.00	\$0		
	TOTAL	OF ABOVE CALCU	LATIONS =	\$890.00		
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	c. [X] The Commissioner is hereby authorized to charge any additional fees which may be required, or credit any overpayment to Deposit Account No. 13-2725.					
	ropriate time limit under 3 filed and granted to resto			petition to revive (37 CFR		
SEND ALL CORRESPONDENCE	то				11111	
Gregory A. Sebald MERCHANT & GOU	JLD	SIG	NATURE:	MUM/		
P.O. Box 2903						
Minneapolis, MN 554	102-0903		NAI	ME: Gregory A. Sebald		
			REG	GISTRATION NUMBER:	33,280	

# IN THE UNITED STATES PATENT AND TRADEMARK OF

Applicant:

**CADELL** 

Docket:

270.63USWO

Title:

METHOD OF CALIBRATING A SPECTROSCOPIC DEVICE

#### **CERTIFICATE UNDER 37 CFR 1.10**

'Express Mail' mailing label number. EV 036304919 US

Date of Deposit February 15, 2002

I hereby certify that this paper or fee is being deposited with the United States Postal Service 'Express Mail Post Office To Addressee' service under 37 CFR 1 10 and is addressed to the Commissioner for Patents, Washington, D.C. 20231

#### BOX PATENT APPLICATION

Commissioner for Patents Washington, D.C. 20231

Sir:

We are transmitting herewith the attached:

- Transmittal sheet, in duplicate, containing Certificate under 37 CFR 1.10.
- National Stage PCT Patent Application: Spec. 15 pgs; 11 claims

The fee has been calculated as shown below in the 'Claims as Filed' table.

- 5 sheets of formal drawings
- An unsigned Combined Declaration and Power of Attorney
- A check in the amount of \$890.00 to cover the Filing Fee
- Other: PTO-1390; Preliminary Amendment with abstract page; Application Data Sheet; Information Disclosure Statement, 1449, cited references; PCT Request; PCT/IB/304; PCT/IB/308; PCT application as published, International Search Report; International Preliminary Examination Report; PCT/IPEA/402
- Return postcard

#### CLAIMS AS FILED

Number of Claims Filed		Iı	n Excess	of:		Numb Extra	er		Rate		Fee
Basic Filing Fee 🗦 👵		2 .	,			_ ,7,			> -		\$890.00
Total Claims		ś.	~ *£	2.		(% <u>.</u> :	4		-;*c		1,12
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TOTAL FILING FEE							1				\$890.00

Please charge any additional fees or credit overpayment to Deposit Account No. 13-2725. A duplicate of this sheet is enclosed.

MERCHANT & GOULD P.C.

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Initials: GAS/tvm

# **Application Data Sheet**

**Application Information** 

Application Type::

Regular

Subject Matter::

Utility

Suggested Classification::

Suggested Group Art Unit::

CD-ROM or CD\_R?::

None

Number of CD disks::

Number of copies of CDs::

Sequence Submission::

No

Computer Readable Form (CRF)?::

No

Title::

METHOD OF CALIBRATING A SPECTROSCOPIC

**DEVICE** 

Attorney Docket Number::

270.63USWO

Request For Early Publication::

No

Request For Non-Publication::

No

Suggested Drawing Figure::

**Total Drawing Sheets::** 

5

Small Entity::

No

Latin Name::

Variety Denomination Name::

Petition Included::

No

Petition Type::

Licensed US Govt. Agency::

**Contract or Grant Numbers::** 

Secrecy Order in Parent Appl.?::

No

## **Applicant Information**

Applicant Authority Type::

Inventor

Primary Citizenship Country::

Canada

Status::

**Full Capacity** 

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Middle Name::

Family Name::

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City of Residence::

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State or Province of Residence::

Ontario

Country of Residence::

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Street of mailing address::

13 Elgin Street East

City of mailing address::

Conestogo

State or Province of mailing address::

Quebec

Country of mailing address::

Canada

Postal or Zip Code of mailing address:: N0B 1N0

## Correspondence Information

Correspondence Customer Number::

23552

# Representative Information

Representative Customer Number::	23552

## **Domestic Priority Information**

Application::	Continuation Type::	Parent Application::	Parent Filing Date::
This is a	National Stage of	PCT/CA00/01005	08/31/00

PCT/CA00/01005	Non-Provisional of	60/151,536	08/31/99

## **Assignee Information**

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Country of mailing address::

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Postal or Zip Code of mailing address:: N2L 5Z4

S/N unknown

**PATENT** 

#### IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

Applicant:

CADELL

Docket No.:

270.63USWO

Serial No.:

unknown

Filed:

concurrent herewith

Int'l Appln No.:

PCT/CA00/01005

Int'l Filing Date:

August 31, 2000

Title:

METHOD OF CALIBRATING A SPECTROSCOPIC DEVICE

CERTIFICATE UNDER 37 CFR 1.10

'Express Mail' mailing label number: EV 036304919 US

Date of Deposit: February 15, 2002

I hereby certify that this correspondence is being deposited with the United States Postal Service 'Express Mail Post Office To Addressee' service under 37 CFR 1.10 on the date indicated above and is addressed to the Assistant Commissioner for Patents, Washington, D.C. 20231.

Name: Chris Stordahl

## PRELIMINARY AMENDMENT

Box PCT Assistant Commissioner for Patents Washington, D.C. 20231

Dear Sir:

In connection with the above-identified application filed herewith, please enter the following preliminary amendment (marked-up copy attached):

#### IN THE ABSTRACT

Insert the attached Abstract page into the application as the last page thereof.

#### IN THE SPECIFICATION

A courtesy copy of the present specification is enclosed herewith. However, the World Intellectual Property Office (WIPO) copy should be relied upon if it is already in the U.S. Patent Office.

IN THE CLAIMS

11. (Amended) A Method according to claim 1, wherein the sample is an

individual patient, the spectroscopic device is a near-infrared spectrophotometer, and the

analyte is selected from the group consisting of glucose, hemoglobin, albumin,

cholesterol, and ethanol.

**REMARKS** 

The above preliminary amendment is made to remove a multiple dependency

from claim 10.

A new abstract page is supplied to conform to that appearing on the publication

page of the WIPO application, but the new Abstract is typed on a separate page as required by

U.S. practice.

Applicants respectfully request that the preliminary amendment described herein

be entered into the record prior to calculation of the filing fee and prior to examination and

consideration of the above-identified application.

If a telephone conference would be helpful in resolving any issues concerning this

communication, please contact Applicants' primary attorney-of record, Gregory A. Sebald (Reg.

No. 33,280), at (612) 336.4728.

Respectfully submitted,

MERCHANT & GOULD P.C.

P.O. Box 2903

Minneapolis, Minnesota 55402-0903

(612) 332-5300

Dated: February 15, 2002

GAS/tvm

Reg. No. 33,280

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## **MARKED-UP COPY**

11. (Amended) A Method according to [any one of claims 1-10] <u>claim 1</u>, wherein the sample is an individual patient, the spectroscopic device is a near-infrared spectrophotometer, and the analyte is selected from the group consisting of glucose, hemoglobin, albumin, cholesterol, and ethanol.

#### IN THE ABSTRACT

Title: METHOD OF CALIBRATING A SPECTROSCOPIC DEVICE

The present invention provides a method of calibrating a spectroscopic device for providing a non-invasive measurement of an analyte level in a sample. The method comprises the steps of: (a) providing a plurality of calibration algorithms; (b) taking a set of non-invasive measurements on said sample with said spectroscopic device; (c) calculating a predicted set of analyte levels for each of the calibration algorithms in response to the set of non-invasive measurements, each of the predicted sets of analyte levels being characterized by a variability range, a slope, an R¿2¿ (a square of the correlation between said set of non-invasive measurements and said predicted set of analyte levels), and a standard error of prediction; and (d) selecting an appropriate calibration algorithm by using a suitability score based on the variability range, the slope, the R² and the standard error of prediction for each of the predicted sets of analyte levels. A method of generating suitable calibration algorithms in step (a) is also provided.

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#### METHOD OF CALIBRATING A SPECTROSCOPIC DEVICE

### FIELD OF THE INVENTION

The present invention relates to the fields of spectroscopy, spectrophotometry, and chemometrics. In particular, the present invention relates to a method of calibrating a spectroscopic device for predicting analyte levels in a non-invasive manner. The method of the present invention is particularly suitable for blood glucose prediction based on near-infrared spectrophotometry measurements.

## **BACKGROUND OF THE INVENTION**

Biotechnological analysis and examination are often based on the measurement of various chemical analytes in the composition of a biological matrix such as blood, interstitial fluid, or living tissue. Such measurements may be used to evaluate a patient's state of health and to determine what, if any, treatment is necessary. For example, the frequent monitoring of blood glucose levels in diabetic persons with glucose meters is often necessary to allow such persons to manage the diabetes mellitus disease, by taking insulin injections or oral drugs to lower blood glucose when required. Intensive treatment based on frequent blood glucose measurements can significantly reduce the incidence of blindness, kidney loss, and other complications associated with diabetes.

Most home-based glucose measurement systems require the patient to invasively collect a blood sample, by pricking his or her finger, placing the sample on an appropriate test strip, and then testing the sample in an optical glucose meter. For millions of diabetics around the world, the use of lancets or other sharp instruments to draw blood for monitoring their insulin levels is a painful process, and one that often builds up calluses on fingers, making the collection of blood even more difficult. This invasive procedure may be especially difficult to perform on children and therefore particularly trying on parents. Furthermore, the test strips required for each blood sample are generally not reusable, and when multiple measurements are taken each day, amount to significant costs from the patient's point of view. Thus, despite the fact that a large number of diabetics should take several measurements throughout each day (for some individuals, physicians recommend

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testing glucose levels from 4 to 7 times daily), due to the pain, cost, and inconvenience involved, many diabetics do not monitor their glucose levels frequently enough.

Spectroscopy is based on the analysis of how incident radiation interacts with the vibrational and rotational states of molecules which are of analytical interest. Spectroscopic measurement techniques have gained increased popularity because of the ability to provide fast and non-invasive measurements of concentrations of different chemicals or analytes. For the reasons indicated above, this is particularly desirable for home based glucose meters. Spectrophotometry is a type of spectroscopy commonly used to quantitatively measure concentrations of analytes based on spectral energy distribution in the absorption spectrum of a sample solution or medium. In spectrophotometry, the energy distribution is typically analyzed within a range of the visible, ultraviolet, infrared, or near-infrared spectra. For example, near-infrared radiation (NIR) is electromagnetic radiation having a wavelength of between about 0.75 and 2.5 micrometers (i.e. from 150 to 400 THz). Near-infrared spectrophotometry generally uses instruments with quartz prisms in monochromators and with lead sulfide photoconductor bands, detectors to observe absorption spectrophotometry is increasingly being used to measure in vivo analytes such as glucose, total hemoglobin, albumin, cholesterol, and ethanol.

Non-invasive, spectrophotometric measurement of glucose in human beings is performed by focusing an incident radiation source (or sources) on a specific part of the body and detecting the spectral distribution of the radiation transmitted therethrough. The absorbance of light from the incident radiation is due to the chemical components within that body part including water, fat, protein, hemoglobin, melanin, and glucose. One difficulty with glucose measurement spectral analysis, is the spectral overlap between glucose and other chemicals found in blood, often in much greater quantities than glucose. In addition, the thickness, color, and structure of the skin, bones, and blood through which the incident radiation passes will affect the transmission (or reflection/absorption) thereof. Furthermore, the concentration of analytes can vary with changes in activity level, diet, hormone

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fluctuations, and other factors. Glucose concentration measurements are also particularly susceptible to variations in physical and chemical conditions including temperature, pressure, humidity, and skin hydration. As a result, to perform a reliable non-invasive glucose prediction, NIR spectral measurements should be performed through a vascular equilibrated region of the body, and a NIR glucose spectrophotometer must be carefully designed so that the quality of raw spectral information from an NIR glucose meter is high. See generally Waynant and Chenault, "Overview of Non-Invasive Optical Glucose Monitoring Techniques", IEEE LEOS Newsletter, vol. 12, no. 2 (April 1998); and Burmeister and Arnold, "Spectroscopic Considerations for Noninvasive Blood Glucose Measurements with Near Infrared Spectroscopy", IEEE LEOS Newsletter, vol. 12, no. 2 (April 1998).

Near-infrared glucose measurements are generally suitable for tissue depths ranging from about 1 mm to 10 cm, and are often performed through a patient's finger tip, although other areas of the body (for example the web between two fingers, an ear lobe, or the upper lip) can also be used. The sample thickness is an important experimental parameter because a greater thickness increases the amount of absorption and thereby lowers the minimum limits for detection, whereas because less incident radiation successfully traverses through a thicker sample (i.e. without being absorbed) effectively increasing the spectral noise: see Burmeister and Arnold, supra.

In addition to the difficulties of obtaining accurate spectral measurements with NIR spectrophotometry, a more significant difficulty associated with such measurements has been the need to calibrate such an instrument so that it may be used by various different individuals, whose analyte concentrations and variability, for example, may differ substantially. Calibration of spectrophotometers (and of analytical instruments in general) is necessary to ensure the accuracy of measurements performed by such devices.

Two approaches to calibrate and predict analyte (e.g. glucose) levels have been used in the prior art. In the first approach, a thorough calibration process is performed separately on each individual with whom the instrument is to be used. This individual calibration process requires taking a number of blood samples over a period of time from

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the individual, and obtaining reference glucose concentration measurements from these samples. A number of corresponding non-invasive spectroscopic measurements are taken concurrently, and calibration regression analysis is then performed to correlate, e.g. through linear regression analysis, the non-invasive measurements with the reference measurements. In this manner, the spectroscopic device or spectrophotometer is custom calibrated to the specific individual. A major disadvantage of this approach, however, is that the calibration model is limited to data from that particular individual which may have an insufficient amount of information regarding variation of spectra overlapping (interfering with) those of the analyte of interest. It is important to be able to characterize such spectra to enable highly accurate analyte predictions. To alleviate this problem, the custom calibration process may have to be carried out frequently, even on a daily basis.

An alternative calibration approach requires that a universal or general calibration algorithm be developed. This approach is based on the notion that by making a large number of calibration measurements (as in the first approach) a single calibration model that sufficiently accounts for all individual variability of all overlapping or interfering analytes can be calculated. Chemometrics, the application of mathematical, statistical and formal logic methods to chemistry, is generally used to process and compute the spectral intensity data and to produce a calibration model: see Small and Arnold, "Data Handling Issues for Near-Infrared Glucose Measurements", IEEE LEOS Newsletter, vol. 12, no. 2 (April 1998); and Shaffer, Small, and Arnold, "Genetic Algorithm-Based Protocol for Coupling Digital Filtering and Partial Least-Squares Regression: Application to the Near-Infrared Analysis of Glucose in Biological Matrices," Anal. Chem. 68, 2663-2675 (1996). However, for measurements of glucose, and other analytes which vary distinctly and differently from one individual sample to another, a single universal calibration algorithm is often ineffective and can result in significant, and sometimes dangerous, erroneous predictions of analyte concentration levels.

Thus, while NIR measurements provide a non-invasive, fast, painless, and convenient technique to monitor glucose levels, correlation

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and clinical interpretation of spectral measurements to obtain the true glucose levels is crucial for proper therapy and disease management. Proper calibration of instruments for different patient populations (which will vary in ethnicity, age, weight, and so on) is crucial in obtaining accurate glucose prediction models. Furthermore, careful validation and testing of the non-invasive results and the glucose prediction equation is needed to determine if the glucose correlation is consistent in all clinically important conditions and for all, or at least most, types of patients.

## **SUMMARY OF THE INVENTION**

In a first aspect the present invention provides a method of calibrating a spectroscopic device for providing a non-invasive measurement of an analyte level in a sample, comprising:

- (a) providing a plurality of calibration algorithms;
- (b) taking a set of non-invasive measurements on said sample with said spectroscopic device;
- (c) calculating a predicted set of analyte levels for each of the calibration algorithms in response to the set of non-invasive measurements, each of the predicted sets of analyte levels being characterized by a variability range, a slope, an R<sup>2</sup> (a square of the correlation between said set of non-invasive measurements and said predicted set of analyte levels), and a standard error of prediction; and
- (d) selecting an appropriate calibration algorithm by using a suitability score based on the variability range, the slope, the R<sup>2</sup> and the standard error of prediction for each of the predicted sets of analyte levels.

Preferably, step (d) comprises a method according to claim 1 wherein, step (d) comprises

- (i) selecting the predicted sets of analyte levels in which the standard error of prediction is less than an upper error limit, the variability range is greater than a lower range limit, and the slope is between a first lower slope limit and an upper slope limit said lower and upper slope limits defining an acceptable slope range;
- (ii) for each of the predicted sets selected in step (i), calculating a suitability score in response to the slope, the  $\mathbb{R}^2$  and the standard error of prediction for that predicted set, and selecting the calibration

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algorithm corresponding to the predicted set having the optimal (highest) suitability score as the appropriate calibration algorithm;

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- (iii) if no predicted sets are selected in step (i), selecting the predicted sets of analyte levels in which the variability range is lower than the lower range limit and in which the standard error of prediction is less than the upper error limit,
- (iv) from each of the predicted sets selected in step (iii), selecting the calibration algorithm corresponding to the predicted set having the lowest standard error of prediction as the appropriate calibration algorithm; and
- (v) if no predicted sets are selected in step (i), or step (iii), determining that no calibration algorithm is appropriate.

Also preferably, the acceptable slope range may be subdivided into a plurality of ubranges corresponding to a plurality of levels, comprising a first level and one or more subsequent levels, and if no predicted sets are selected in the first level, then repeating step (i) at each subsequent level until a predicted set is selected or there are no more subsequent levels.

Preferably, the plurality of calibration algorithms provided in step (a) are first generated by the steps of:

- (i) compiling non-invasive and corresponding reference data sets of measurements of analyte levels for each of a number of samples;
  - (ii) rejecting data sets that are not suitable for calibration;
- (iii) combining data sets that are suitable for calibration into a plurality of groups depending on whether correlations of the combined data sets meet predetermined criteria; and
- (iv) generating a calibration algorithm for each of the groups of data sets.

Steps (a), (c), and (d) of the method of the present invention may be performed on a computer associated with the spectroscopic device. Furthermore, the sample may be an individual patient, the spectroscopic device may be a near-infrared spectrophotometer, and the analyte may be selected from the group consisting of glucose, hemoglobin, albumin, cholesterol, and ethanol.

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The objects and advantages of the present invention will be better understood and more readily apparent with reference to the remainder of the description in conjunction with the accompanying drawings.

## **BRIEF DESCRIPTION OF THE DRAWINGS**

In the drawings which illustrate, by way of example, a preferred embodiment of the invention:

Figure 1 is a correlation scatterplot of glucose predictions for a number of different patients based on a universal, single calibration model;

Figure 2 is a correlation scatterplot of glucose predictions for a number of patients based on the multiple algorithm calibration model according to the present invention;

Figure 3 is another correlation scatterplot of glucose predictions for a number of patients based on the multiple algorithm calibration model according to the present invention;

Figure 4 is a correlation scatterplot of glucose predictions for a typical patient based on a universal, single calibration model; and

Figure 5 is a correlation scatterplot of glucose predictions for the same typical patient based on the multiple algorithm calibration model according to the present invention.

## **DETAILED DESCRIPTION OF THE INVENTION**

As mentioned, the present invention relates to a method of calibrating a spectroscopic device for predicting analyte levels in a non-invasive manner. The present invention can be used with a typical NIR spectrophotometer system having a light source which is projected through the item to be examined, a sample interface mechanism, a spectrometer to separate the light into its component wavelengths, a detector, amplification electronics and a computer. By measuring the loss (absorption), between the source and the detector and applying appropriate chemometric (mathematical) techniques, it is possible to non-invasively determine the chemicals being examined since different chemicals absorb different amounts of light at different wavelengths. Such a spectrophotometric device and method are described in detail in United States Patent No. 5,361,758, the contents of which are incorporated herein by virtue of this reference. While the present description relates primarily to glucose measurement, one of the major

fields of application for NIR measurement at present, it will be understood that the principles of the present invention equivalently apply to other analytes non-invasively measured using various spectroscopic techniques.

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To utilize the NIR spectrum for glucose measurement, it is necessary to use a spectrometer which has wide dynamic range, a high signal to noise ratio, and exhibits low scattering losses. The output from the spectrometer is used to generate spectra with high precision both in absorbance and wavelength. Significant glucose absorption bands are centered about wavelengths of 1.67, 2.13, 2.27, and 2.33 micrometers (as discussed in Small and Arnold, "Data handling Issues for Near-Infrared Glucose Measurements", supra). In addition to these wavelengths there are significant glucose absorption bands at 0.97 and 1.12 micrometers which wavelengths allow transmission measurement through a greater distance in the tissue. To be able to use NIR to measure a particular compound/analyte, chemometric mathematical analysis is applied to the measured spectrum. The mathematical analysis techniques are carried out by a computer equipped with advanced software capable of interpreting the resulting complex spectra.

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To universally calibrate an NIR spectrophotometer, the absorbance associated with the compound or analyte of interest must first be measured on a relatively large number of samples. These NIR measurements are then compared to measurements made in a more traditional and more accurate invasive manner. From these comparisons an algorithm is developed that characterizes the analyte to be measured. The methods used to generate calibration models for NIR spectrophotometers are sophisticated. An important criterion is that calibration samples be uncorrelated with other chemical species concentrations. Satisfaction of these criteria is burdensome because of the large number of chemical species typically present in a biological matrix or growth medium.

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The present invention provides a multi-algorithm calibration method for use with a spectroscopic device capable of predicting analyte levels in a biological matrix, such as blood glucose levels, for a variety of different samples with significantly greater accuracy than prior art universal single calibration models. The calibration method of the

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invention has two key parts: the generation of a plurality of different algorithms and the selection of the algorithm which is most appropriate for generating an accurate prediction of analyte concentration.

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To generate suitable calibration algorithms, the non-invasive and reference measurement data sets for a number of individuals are first compiled. Once this is achieved, correlation techniques are carried out to relate non-invasive spectral measurements to reference glucose levels. At this stage the suitability of individual data samples for calibration purposes must be considered. If some of the non-invasive spectral measurements in a data set result in a saturation of the measurement equipment, e.g. at the output of an analog-to-digital converter, then the data sample should be disqualified for use in a calibration algorithm. Similarly, if the range of variability of the non-invasive spectral measurements in a data set is too small, the data sample should also be disqualified from calibration. Furthermore, if the correlation of glucose in a particular data set is undesirably correlated to another variable, such as the concentration of another chemical, the sample should again be disqualified. As a result, a large numbers of sample sets will not be suitable for calibration algorithm development.

The grouping or combining of calibration-suitable data sets (each specific to a particular individual) into different algorithms occurs on a trial and error basis, by repeatedly combining these data sets and performing a new correlation on the combined sets of data. If the correlations of the combined sets meet certain criteria, then the combination is acceptable, if not, a different combination is attempted. Once a combination is deemed acceptable, the addition of a further data set is attempted and its acceptability is determined in turn. In this manner, a combination of data sets for an algorithm grows from 2 data sets, to 3 data sets, to 4 data sets, and so on until a sufficiently large number of data sets is obtained from which a reliable calibration algorithm can be developed.

In combining data sets for a single algorithm, two criteria are key. First, any correlation of glucose in the combined data set group with other parameters must be minimized. Such undesirable correlations may be exhibited with different chemical species concentrations or other time varying quantities. A suitable quantitative test for this criteria is to

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ensure that all undesirable correlations of the combined data sets, as indicated by the square of the correlation coefficient  $R_{\rm u}^2$  (for the undesirable correlations), are less than a certain limit, such as  $R_{\rm u}^2 < 0.10$ . In known manner, the correlation coefficient,  $R_{\rm u}$ , is a measure of the tendency of two variables to vary together. If the addition of a data set to another data set or to a suitable sub-combination of data sets does not satisfy this criteria, the new combination will not be acceptable.

In addition, the combination of data sets can be optimized by ensuring that the data sets exhibit a high degree of correlation with one another, so a second criteria in combining data sets should require that the data sets in a particular group be sufficiently correlated with one another, This can be achieved, for example, by ensuring that the Standard Error of Prediction (SEP) is minimal and the square of the correlation coefficient, R<sup>2</sup>, together with the slope relating reference glucose to predicted glucose, are greater than about 0.8 for an existing sub-combination of data sets and another data set being potentially combined with that sub-combination.

In an alternate embodiment of the present invention, instead of successively adding on to, or building up, groups of data sets for a calibration algorithm, a large group of calibration suitable data sets can be initially taken up at random. In this embodiment, data sets are removed from the group, and the SEP and correlation criteria above are reevaluated. If the SEP, slope and correlation criteria improve after the removal of the data set, the data set is permanently removed from the group, whereas if the criteria do not improve, the data set is reinstated back into the group. Using this top-down approach, the removal of data sets can cease when the criteria meet acceptable limits.

The SEP, slope, and correlation calculations for determination of the calibration algorithm groupings (and the associated calibration algorithms) become increasingly complex with increasing amounts of data and are preferably carried out using partial least-squares (PLS) regression analysis. The PLS analysis techniques are well known to those skilled in the art and provide good linear approximation by removing considerable redundant information: see generally Small and Arnold, "Data handling Issues for Near-Infrared Glucose Measurements", supra.

Other types of analysis, such as partial component analysis (PCA) or artificial neural networks (ANN), can also be used.

This calibration algorithm generation process is used to generate at least two groups of data sets (corresponding to groups of individuals), each providing a calibration model or algorithm for predicting the level of analyte present in a biological matrix based on a non-invasive spectroscopic measurement. Preferably, two initial groups of data set are developed, and, from these two groups, additional groups can be generated. These additional groups are generally smaller in size than the two initial groups and may meet more stringent correlation criteria. These additional groups may be constructed with the top-down approach described above, wherein data sets are selectively removed from an initial group to determine if the correlation criteria improve or deteriorate after the removal.

Given the availability of multiple calibration algorithms, it is necessary to select the most suitable of these algorithms for predicting the levels of analyte in any given patient or sample. Once this is done the spectroscopic device is calibrated with the selected algorithm. The algorithm selection process is generally independent of the process used to generate the multiple algorithms, although it will clearly depend on the number of different algorithms that are available.

For any given patient or sample on which the non-invasive spectroscopic measurement is to be made, it is necessary first to compile a set of highly accurate reference measurements and corresponding non-invasive spectral measurements at selected intervals over a period of time. In a preferred embodiment, two successive non-invasive spectral measurements are successively taken each measurement interval, and their mean is then taken as a single non-invasive measurement for that interval.

Before algorithm selection begins, two variables may be set to determine the stringency of acceptance requirements for patients (i.e. whether at least one of the available calibration algorithms will be found suitable for a patient). These are (i) a lower range limit for the range of glucose or analyte variability in the data set (if the data does not vary significantly it may not provide sufficient information to be useful), and (ii) an upper error limit (SEP<sub>max</sub>) for the standard error of prediction

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(SEP), i.e. the square root of the mean of the squared deviations of the reference analyte concentrations from the analyte concentration values predicted by a calibration algorithm.

From the non-invasive spectral measurements, a predicted set of analyte concentration levels are calculated according to each calibration algorithm. When compared to the reference set of concentration levels, each predicted set will be calculated from a linear regression or "best fit" line characterized by: a range of glucose level variability; a slope (which ideally equals 1, i.e. the predicted values correspond directly to the reference values); a correlation coefficient R, defining the correlation of the predicted value set with the reference value set; and a standard error of prediction (SEP), as defined above. Note that as the correlation coefficient R increases the SEP will decrease, and vice versa, except in cases in which there is an offset or bias in the predicted analyte.

By way of example, and not to restrict the scope of the present invention, the algorithm selection process can include 4 levels or steps. In a first level, if the predicted data set for an algorithm meets the following criteria

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- i) SEP < upper error limit (SEP<sub>max</sub>)
- ii) Analyte range > lower range limit

and iii) 0.5 < slope (of regression line) < 1.05,

a suitability score is calculated according to the following equation:

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$$score = (slope) (R^2) / (SEP)$$

If more than one algorithm meets the first level criteria, the algorithm with the highest suitability score is selected and the selection process is complete. If no algorithm qualifies in terms of the first level criteria, the selection process advances to a second level.

The second level is identical to the first level, except that the slope criteria becomes

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0.33 < slope (of regression line) < 0.5

For algorithms that meet this revised slope criterion (and the SEP and range criteria), a suitability score is calculated as in the first level. Again, if more than one algorithm qualifies at the second level, the algorithm with the highest score is selected, and the selection process is complete.

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At a third level, if none of the algorithms qualify at either the first level or the second level and the range of analyte measurements is greater than the lower range limit, the sample or patient is excluded from measurement, since under these circumstances no calibration algorithm can be relied on to perform accurately.

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Lastly, at a fourth level, if none of the algorithms qualify at either the first level or the second level and the range of analyte measurements is lower than the lower range limit, the algorithm with the lowest SEP is selected if that SEP is less than the upper error limit. If the SEP is greater than the upper error limit, the patient is again excluded from measurement. Thus, in the fourth level, the SEP becomes the sole factor in deciding whether an algorithm should be accepted.

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Note that the slope criteria limits in the first and second levels described above are preferred limits, and these levels can also be varied somewhat (similar to the lower range limit and upper error limit) depending on the application and acceptance requirements. Also, the number of levels or steps used in the algorithm selection process can vary, from a single level covering the entire preferred range, to two or more levels subdividing the preferred range into two or more range limits or range criterion. Regardless of the number of levels, however, the third and fourth levels as described above will correspond to the last two levels, in which the patient is either excluded, or a decision to accept an algorithm is made on the basis of SEP alone. It will be readily appreciated by those skilled in the art that the number of levels or steps chosen is a matter of preference only.

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To illustrate the principals of the present invention, the multiple algorithm calibration method was used to calibrate a NIR spectrophotometer for measuring glucose levels in a number of different patients. All reference data measurements were taken with a glucose analyzer such as the industry standard YSI<sup>TM</sup> (Yellow Springs Instruments) Glucose Analyzer which provides precise glucose measurements on (invasive) blood samples. The inventor found that

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only approximately 6% of single NIR patient data sets qualified for calibration purposes, given the criteria described above. These calibration-qualified sets of data were grouped into two initial calibration algorithm sets, each initial group was built up to a group with 33 sets of data within it. Eleven additional groups were spun off from these two initial groups and each of these groups had at least 8 sets of data within it. Thus, thirteen different calibration algorithms were available for predicting glucose levels.

The algorithm selection was based on a total of 24 reference measurements (i.e. YSI<sup>TM</sup> measured) and 24 means of two NIR finger measurements, taken from a patient over a three day period. The above described algorithm selection process was run for a number of different patients (not involved in the calibration process).

Figure 1 shows a correlation scatterplot of glucose predictions for a number of different patients based on a universal calibration model (QUAD 37). The scatterplot in Figure 1 is superimposed on a Clarke error grid, as described in Clarke and Cox, "Error Grid Analysis", Diabetes Care, 10:622-628 (1987). The Clarke error grid breaks the correlation space into five regions (A-E) that assess measurement accuracy on the basis of validity of the corresponding clinical decision. Correlation points falling within the "A" region correspond to the correct clinical decision being made based on the similarity between the actual and predicted glucose levels. (i.e. the predicted values deviate by no more than 20% from the reference values). In the "B" region, the predicted values deviate by more than 20% from the reference values, but treatment decisions made based on the predicted levels of glucose would not jeopardize or adversely affect the patient. In regions "C", "D", and "E", the predictions significantly deviate from the reference values, and treatment decisions based on these predictions may well be harmful to a patient. The Clarke error grid is often used to evaluate the clinical consequences of home-use blood glucose monitor errors, in contrast to more conventional analytical methods which may be more suitable for laboratory reference devices.

Referring to Figure 1, it can be seen that the upper error limit SEP<sub>max</sub> for this universal calibration algorithm is 3.07 mmol/L and that significant amounts of the data are outside acceptable limits or regions.

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In contrast, Figure 2 shows the correlation scatterplot of glucose predictions for 25 patients based on the multiple algorithm calibration model according to the present invention, with SEP<sub>max</sub> equal to 2.76 mmol/L. Similarly, Figure 3 shows a correlation scatterplot of glucose predictions for 22 patients based on the multiple algorithm calibration model according to the present invention, with SEP<sub>max</sub> equal to 2.28 mmol/L. Both the predictions in Figure 2 and particularly Figure 3 exhibit much better accuracy than those in Figure 1 (calibrated according to a universal algorithm).

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Similarly, Figure 4 shows a set of correlation data for a typical patient, calibrated according to the same universal algorithm, whereas Figure 5 shows the correlation data for the identical patient calibrated according to the multiple algorithm method of the present invention. Once more, the improved accuracy of the glucose predictions based on the calibration method of the present invention is evident.

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It will be clear to those skilled in the art that the principles of the present invention are applicable to a broad range of spectroscopic applications. For example, NIR radiation can be used to penetrate such items as human tissue, vials of blood, or containers of milk - all normally considered opaque. Mathematical analysis of the resulting absorption spectrum determines the composition of the substance penetrated by the light. Thus, in a blood testing lab, samples can be examined without contact by passing the light through a plastic vial containing the liquid. Similarly, in a dairy application, the NIR light is passed through a bag of milk and such parameters as butterfat, solid particles and lactate are measured. The multiple algorithm method of the present invention is easily extendible to these types of non-invasive measurements as well. Furthermore, the method of the present invention can also be applied to the non invasive measurement of a number of other blood analytes such as cholesterol, hemoglobin, Hb1Ac, Fructosamine, and 1.5 AG.

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While preferred embodiments of the present invention have been described, the embodiments disclosed are illustrative and not restrictive, and the invention is intended to be defined by the appended claims.

## I CLAIM:

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- 1. A method of calibrating a spectroscopic device for providing a non-invasive measurement of an analyte level in a sample, comprising:
  - (a) providing a plurality of calibration algorithms;
- (b) taking a set of non-invasive measurements on said sample with said spectroscopic device;
- (c) calculating a predicted set of analyte levels for each of the calibration algorithms in response to the set of non-invasive measurements, each of the predicted sets of analyte levels being characterized by a variability range, a slope, an R<sup>2</sup> (a square of the correlation between said set of non-invasive measurements and said predicted set of analyte levels), and a standard error of prediction; and
- (d) selecting an appropriate calibration algorithm by using a suitability score based on the variability range, the slope, the R<sup>2</sup> and the standard error of prediction for each of the predicted sets of analyte levels.
- 2. A method according to claim 1 wherein, step (d) comprises
- (i) selecting the predicted sets of analyte levels in which the standard error of prediction is less than an upper error limit, the variability range is greater than a lower range limit, and the slope is between a first lower slope limit and an upper slope limit said lower and upper slope limits defining an acceptable slope range;
- (ii) for each of the predicted sets selected in step (i), calculating a suitability score in response to the slope, the R<sup>2</sup> and the standard error of prediction for that predicted set, and selecting the calibration algorithm corresponding to the predicted set having the optimal (highest) suitability score as the appropriate calibration algorithm;
- (iii) if no predicted sets are selected in step (i), selecting the predicted sets of analyte levels in which the variability range is lower than the lower range limit and in which the standard error of prediction is less than the upper error limit,
- (iv) from each of the predicted sets selected in step (iii), selecting the calibration algorithm corresponding to the predicted set having the

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lowest standard error of prediction as the appropriate calibration algorithm; and

- (v) if no predicted sets are selected in step (i), or step (iii), determining that no calibration algorithm is appropriate.
- 3. A method according to claim 2 wherein, said acceptable slope range is subdivided into a plurality of subranges corresponding to a plurality of levels, comprising a first level and one or more subsequent levels, and if no predicted sets are selected in the first level, then repeating step (i) at each subsequent level until a predicted set is selected or there are no more subsequent levels.
- 4. A method according to claim 2 wherein, said lower slope limit is less than one and said upper slope limit is greater than one.
- 5. A method according to claim 4 wherein, said lower slope limit is about 0.3 and said upper slope limit is about 1.05.
- 6. A method according to claim 3, wherein, said acceptable slope range is defined by a lower slope limit of about 0.3 an upper slope limit of about 1.05.
  - 7. A method according to claim 1 wherein, the plurality of calibration algorithms provided in step (a) are first generated by the steps of:
  - (i) compiling non-invasive and corresponding reference data sets of measurements of analyte levels for each of a number of samples;
    - (ii) rejecting data sets that are not suitable for calibration;
  - (iii) combining data sets that are suitable for calibration into a plurality of groups depending on whether correlations of the combined data sets meet predetermined criteria; and
  - (iv) generating a calibration algorithm for each of the groups of data sets.
- 8. A method according to claim 7 wherein the predetermined criteria in step (iii) include minimizing correlations of the combined data

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sets in a particular group with parameters other than said analyte and maximizing the correlation between data sets in a particular group.

- 9. A method according to claim 7 wherein steps (iii) and (iv) are performed using partial least-squares regression analysis.
- 10. A method according to claim 7 wherein steps (a), (c), and (d) are performed on a computer associated with the spectroscopic device.
- 10 11. A method according to any one of claims 1-10 wherein the sample is an individual patient, the spectroscopic device is a near-infrared spectrophotometer, and the analyte is selected from the group consisting of glucose, hemoglobin, albumin, cholesterol, and ethanol.

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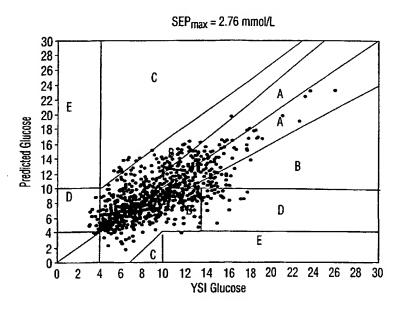
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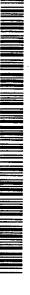
#### (54) Title: METHOD OF CALIBRATING A SPECTROSCOPIC DEVICE



Clarke Error Grid

% Zone 52.24 Α В 43.37 C 1.07 Ď 3.22 0.10

(57) Abstract: The present invention provides a method of calibrating a spectroscopic device for providing a non-invasive measurement of an analyte level in a sample. The method comprises the steps of: (a) providing a plurality of calibration algorithms; (b) taking a set of non-invasive measurements on said sample with said spectroscopic device; (c) calculating a predicted set of analyte levels for each of the calibration algorithms in response to the set of non-invasive measurements, each of the predicted sets of analyte levels being characterized by a variability range, a slope, an R¿2¿ (a square of the correlation between said set of non-invasive measurements and said predicted set of analyte levels), and a standard error of prediction; and (d) selecting an appropriate calibration algorithm by using a suitability score based on the variability range, the slope, the R2 and the standard error of prediction for each of the predicted sets of analyte levels. A method of generating suitable calibration algorithms in step (a) is also provided.



WO 01/16579 AJ

10/049699

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Docket No . 270 63USWO
Title. METHOD OF CALIBRATING A SPECTROSCOPIC DEVICE
Attorney Name Gregory A. Sebald
Phone No.. 612.336 4728
Sheet 1 of 5

 $SEP_{max} = 3.07 \text{ mmol/L}$ 

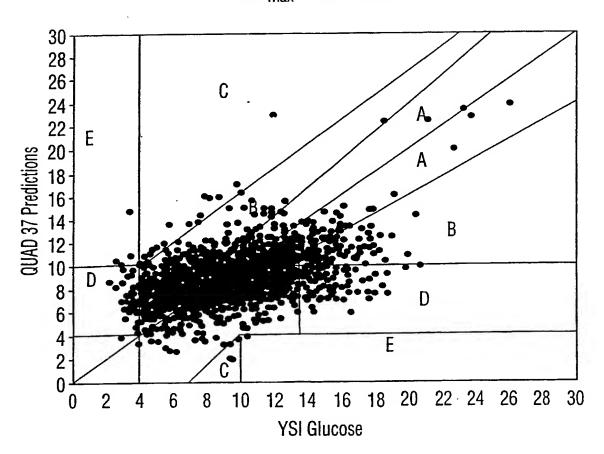
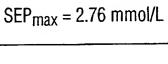
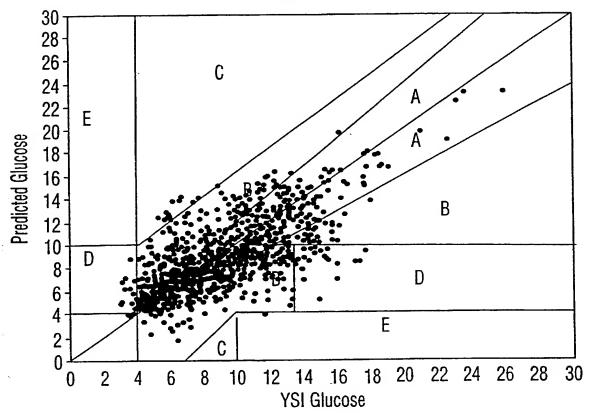


FIGURE 1

Inventor CADELL Docket No 270.63USWO Title. METHOD OF CALIBRATING A SPECTROSCOPIC DEVICE Attorney Name Gregory A Sebald Phone No 612.336 4728 Sheet 2 of 5





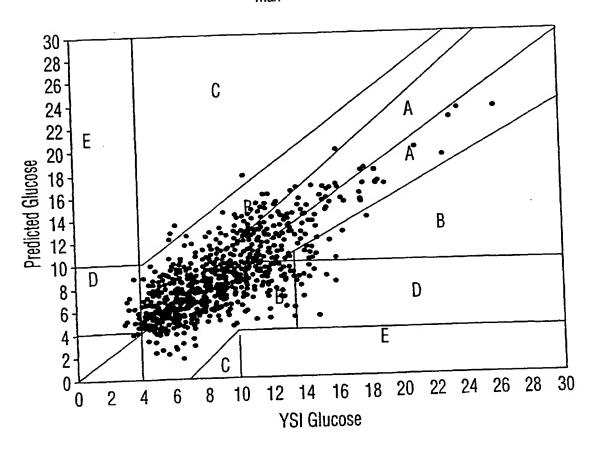
# Clarke Error Grid

Zone	%
Α	52.24
В	43.37
C	1.07
D	3.22
Ε	0.10

# FIGURE 2

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Sheet 3 of 5

 $SEP_{max} = 2.28 \text{ mmol/L}$ 



# Clarke Error Grid

Zone	%
Α	54.11
В	43.31
C	0.49
Ď	1.96
Ε	0.12

# FIGURE 3

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Title METHOD OF CALIBRATING A SPECTROSCOPIC DEVICE
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Phone No 612.336 4728
Sheet 4 of 5

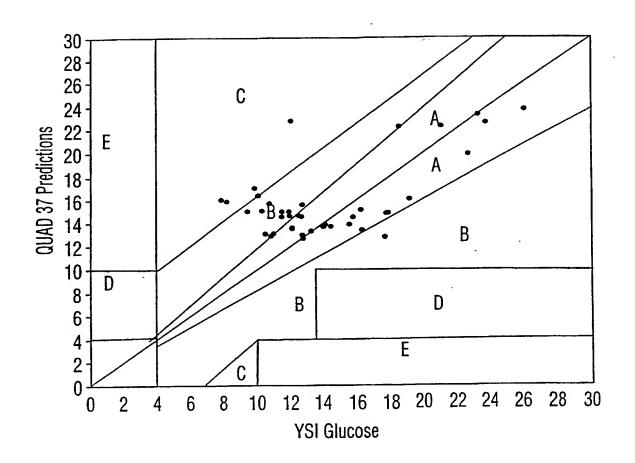


FIGURE 4

10/049699

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Title METHOD OF CALIBRATING A SPECTROSCOPIC DEVICE
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Phone No · 612.336 4728
Sheet 5 of 5

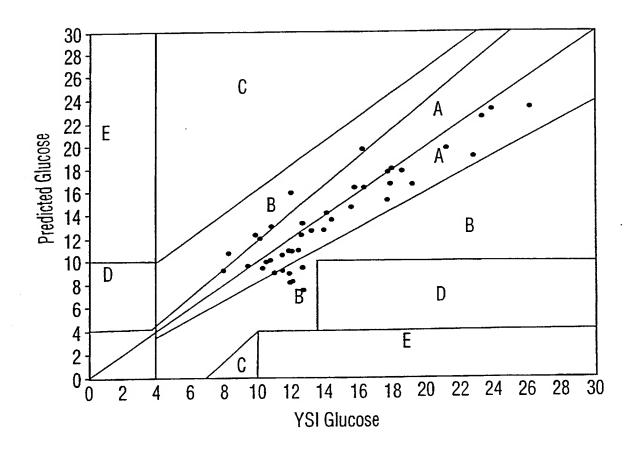


FIGURE 5

The specification of which 1. is attached hereto

1: 1

CME TELEMETRIX

Anomes Docker No. 270.63USWO

RCHANT & GOULD P.C. ted States Patent Application Q0/049022

## COMBINED DECLARATION AND POWER OF ATTORNEY

As a below named inventor I hereby declare that: my residence, post office address and citizenship are as stated below next to my iame; that

I verily believe I am the original, first and sole inventor (if only one name is listed below) or a joint inventor (if plural inventors are named below) of the subject matter which is claimed and for which a patent is sought on the invention entitled: METHOD OF CALIBRATING A SPECTROSCOPIC DEVICE

<ol> <li>was filed on as described and claimed in and for which I solicit a U</li> </ol>	application serial no. — and was amer international no. PCT/CA00/01005 file Juited States patent.	d August 31, 2000 and as ame	nded on (if any), which I have review	ved
I hereby state that I have any amendment referred		f the above-identified specific	ation, including the claims, as amended t	у
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	FOREIGN APPLICATION(S), IF ANY	, CLAIMING PRIORITY UNDER	35 USC § 119	=
COUNTRY	APPLICATION NUMBER	DATE OF FILING	DATE OF ISSUE	
		(day, month, year)	(day, month, year)	1
				1
	all foreign application(s), if any,	filed before the priority	APPLICATION(S)	_
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		(day, month, year)	(day, month, year)	4

I hereby claim the benefit under Title 35, United States Code, § 120/365 of any United States and PCT international application(s) listed below and, insofar as the subject matter of each of the claims of this application is not disclosed in the prior United States application in the manner provided by the first paragraph of Title 35, United States Code, § 112, I acknowledge the duty to disclose material information as defined in Title 37, Code of Federal Regulations, § 1.56(a) which occurred between the filing date of the prior application and the national or PCT international filing date of this application.

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# CME TELEMETRIX 10049699, 161906

acknowledge the duty to disclose information that is material to the patentability of this application in accordance with Title 37, Code of ederal Regulations, § 1.56 (reprinted below):

## 1.56 Duty to disclose information material to patentability.

- (a) A patent by its very nature is affected with a public interest. The public interest is best served, and the most effective atent examination occurs when, at the time an application is being examined, the Office is aware of and evaluates the teachings of all information material to patentability. Each individual associated with the filing and prosecution of a patent application has a duty of candor and good faith in dealing with the Office, which includes a duty to disclose to the Office all information known to that individual to be naterial to patentability as defined in this section. The duty to disclose information exists with respect to each pending claim until the claim is canceled or withdrawn from consideration, or the application becomes abandoned. Information material to the patentability of a claim that is canceled or withdrawn from consideration need not be submitted if the information is not material to the patentability of any claim remaining under consideration in the application. There is no duty to submit information which is not material to the patentability of my existing claim. The duty to disclose all information known to be material to patentability is deemed to be satisfied if all information mown to be material to patentability of any claim issued in a patent was cited by the Office or submitted to the Office in the manner prescribed by §§ 1.97(b)-(d) and 1.98. However, no patent will be granted on an application in connection with which fraud on the Office was practiced or attempted or the duty of disclosure was violated through bad faith or intentional misconduct. The Office encourages applicants to carefully examine:
  - (1) prior art cited in search reports of a foreign patent office in a counterpart application, and
- (2) the closest information over which individuals associated with the filing or prosecution of a patent application believe any pending claim patentably defines, to make sure that any material information contained therein is disclosed to the Office.
- (b) Under this section, information is material to patentability when it is not cumulative to information already of record or being made of record in the application, and
  - (1) It establishes, by itself or in combination with other information, a prima facie case of unpatentability of a claim;
  - (2) It refutes, or is inconsistent with, a position the applicant takes in:
    - (i) Opposing an argument of unpatentability relied on by the Office, or
    - (ii) Asserting an argument of patentability.

A prima facie case of unpatentability is established when the information compels a conclusion that a claim is unpatentable under the preponderance of evidence, burden-of-proof standard, giving each term in the claim its broadest reasonable construction consistent with the specification, and before any consideration is given to evidence which may be submitted in an attempt to establish a contrary conclusion of patentability.

- (c) Individuals associated with the filing or prosecution of a patent application within the meaning of this section are:
  - (1) Each inventor named in the application:
  - (2) Each attorney or agent who prepares or prosecutes the application; and
- (3) Every other person who is substantively involved in the preparation or prosecution of the application and who is associated with the inventor, with the assignee or with anyone to whom there is an obligation to assign the application.
- (d) Individuals other than the attorney, agent or inventor may comply with this section by disclosing information to the attorney, agent, or inventor.
- (e) In any continuation-in-part application, the duty under this section includes the duty to disclose to the Office all information known to the person to be material to patentability, as defined in paragraph (b) of this section, which became available between the filing date of the prior application and the national or PCT international filing date of the continuation-in-part application.

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3umham, Jonelle	Reg. No. 41,980	Pauly, Daniel M.	Reg. No. 40,123
Byrne, Linda M.	Reg. No 32,404	Phillips, John B.	Reg. No. 37,206
Campbell, Keith	Reg. No. 46,597	Pino, Mark J.	Reg. No. 43,858
	Reg. No. 25,959	Prendergast, Paul	Reg. No. 46,068
Carlson, Alan G.	Reg. No. 33,227	Pytel, Melissa J.	Reg. No. 41,513
Caspers, Philip P.	Reg. No. 30,247	Qualey, Terry	Reg. No. 25,148
Clifford, John A.	Reg. No. 48,649	Randall, Joshua N.	Reg. No. P-50,719
Cook, Jeffrey	Reg. No. 25,968	Reich, John C.	Reg. No. 37,703
Daignault, Ronald A.		Reiland, Earl D.	Reg. No. <del>25,7</del> 67
Daley, Dennis R.	Reg. No. 34,994	Schmaltz, David G.	Reg. No. 39,828
Daulton, Julie R.	Reg. No. <u>36,4</u> 14 Reg. No. <u>42,15</u> 7	Schuman, Mark D.	Reg. No. 31,197
DeVries Smith, Katherine M.		· ·	Reg. No. 30,422
DiPietro, Mark J.	Reg. No. 28,707	Schumann, Michael D.	Reg. No. 42,137
Doscotch, Matthew A.	Reg. No. 48,957	Scull, Timothy B.	Reg. No. 33,280
Edell, Robert T.	Reg. No. 20,187	Sebald, Gregory A. Skoog, Mark T.	
Epp Ryan, Sandra	Reg. No. 39,667		Reg. No. 40,178
Fitzsimmons, Karen A.	Reg. No. P-50,470	Spellman, Steven J.	Reg. No. 45,124
Goff, Jared S.	Reg. No. 44,716	Stewart, Alan R.	Reg. No. 47,974
Goggin, Matthew J.	Reg. No. 44,125	Stoll-DeBell, Kirstin L.	Reg. No. 43,164
Golla, Charles E.	Reg. No. <del>26,</del> 896	Sullivan, Timothy	Reg. No. 47,981
Gorman, Alan G.	Reg. No. $38,472$	Sumner, John P.	Reg. No. 29,114
Gould, John D.	Reg. No. 18,223	Swenson, Erik G.	Reg. No. 45,147
Gregson, Richard	Reg. No. 41,804	Tellekson, David K.	Reg. No. 32,314
Gresens, John J.	Reg. No. 33,112	Trembath, Jon R.	Reg. No. 38,344
Hamer, Samuel A.	Reg. No. 46,754	Tunheim, Marcia A.	Reg. No. 42,189
Hamre, Curtis B.	Reg. No. 29,165	Underhill, Albert L.	Reg. No. 27,403
Harrison, Kevin C.	Reg. No. 46,759	Vandenburgh, J. Derek	Reg. No. 32,179
Hennings, Mark	Reg. No. 48,982	Wahl, John R.	Reg. No. 33,044
Hertzberg, Brett A.	Reg. No. 42,660	Weaver, Paul L.	Reg. No. 48,640
Hillson, Randall A.	Reg. No. 31,838	Welter, Paul A.	Reg. No. 20,890
Holzer, Jr., Richard J.	Reg. No. 42,668	Whipps, Brian	Reg. No. 43,261
Hope, Leonard J.	Reg. No. 44,774	Whitaker, John E.	Reg. No. 42,222
Jardine, John S.	Reg. No. 48,835	Wier, David D.	Reg. No. 48,229
Johns, Nicholas P.	Reg. No. 48,995	Williams, Douglas J.	Reg. No. 2 <del>7,05</del> 4
Johnston, Scott W.	Reg. No. 39,721	Withers, James D.	Reg. No. 40.376
Kadievitch, Natalie D.	Reg. No. 34,196	Wong, Bryan	Reg. No. P-50,836
Kalinsky, Robert A.	Reg. No. P-50,471	Wong, Thomas S.	Reg. No. 48,577
Kettelberger, Denise	Reg. No. 33,924	Young, Thomas	Reg. No. 25,796
Keys, Jeramie J.	Reg. No. 42,724	Zeuli, Anthony R.	Reg. No. 45.255
Knearl, Homer L.	Reg. No. 21,197		
Kowalchyk, Alan W.	Reg. No. 31,535	•	
Kowalchyk, Katherine M.	Reg. No. 36,848		
Lamberty, Michael	Reg. No. P-50,760		
Lacy, Paul E.	Reg. No. 38,946		
Larson, James A.	Reg. No. 40,443		
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I hereby authorize them to act and rely on instructions from and communicate directly with the person/assignee/attorney/firm/ organization who/which first sends/sent this case to them and by whom/which I hereby declare that I have consented after full disclosure to be represented unless/until I instruct Merchant & Gould P.C. to the contrary.

I understand that the execution of this document, and the grant of a power of attorney, does not in itself establish an attorney-client relationship between the undersigned and the law firm Merchant & Gould P.C., or any of its attorneys.

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hereby declare that all statements made herein of my own knowledge are true and that all statements made on information and belief are relieved to be true; and further that these statements were made with the knowledge that willful false statements and the like so made are sunishable by fine or imprisonment, or both, under Section 1001 of Title 18 of the United States Code and that such willful false statements nay jeopardize the validity of the application or any patent issued thereon.

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